

# SEQUENCE LISTING

<110> Bristol-Myers Squibb Company

<120> POLYNUCLEOTIDE ENCODING A NOVEL HUMAN SERPIN SECRETED FROM LYMPHOID CELLS, LSI-01

<130> D0051.NP

<150> US 60/248,434

<151> 2000-11-14

<150> US 60/257,610

<151> 2000-12-21

<150> US 60/282,745

<151> 2001-04-10

<160> 46

<170> PatentIn version 3.0

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<212> DNA

<213> Homo sapiens

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Phe Cys Ser Lys Met Ala Ser Tyr Leu Tyr Gly Val Leu Phe Ala Val  
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Gly Leu Cys Ala Pro Ile Tyr Cys Val Ser Pro Ala Asn Ala Pro Ser  
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gca tac ccc cgc cct tcc tcc aca aag agc acc cct gcc tca cag gtg 253  
Ala Tyr Pro Arg Pro Ser Ser Thr Lys Ser Thr Pro Ala Ser Gln Val  
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Tyr Ser Leu Asn Thr Asp Phe Ala Phe Arg Leu Tyr Arg Arg Leu Val  
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Leu Glu Thr Pro Ser Gln Asn Ile Phe Phe Ser Pro Val Ser Val Ser  
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Thr Ser Leu Ala Met Leu Ser Leu Gly Ala His Ser Val Thr Lys Thr	
95 100 105 110	
cag att ctc cag ggc ctg ggc ttc aac ctc aca cac aca cca gag tct	445
Gln Ile Leu Gln Gly Leu Gly Phe Asn Leu Thr His Thr Pro Glu Ser	
115 120 125	
gcc atc cac cag ggc ttc cag cac ctg gtt cac tca ctg act gtt ccc	493
Ala Ile His Gln Gly Phe Gln His Leu Val His Ser Leu Thr Val Pro	
130 135 140	
agc aaa gac ctg acc ttg aag atg gga agt gcc ctc ttc gtc aag aag	541
Ser Lys Asp Leu Thr Leu Lys Met Gly Ser Ala Leu Phe Val Lys Lys	
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Glu Leu Gln Leu Gln Ala Asn Phe Leu Gly Asn Val Lys Arg Leu Tyr	
160 165 170	
gaa gca gaa gtc ttt tct aca gat ttc tcc aac ccc tcc att gcc cag	637
Glu Ala Glu Val Phe Ser Thr Asp Phe Ser Asn Pro Ser Ile Ala Gln	
175 180 185 190	
gcg agg atc aac agc cat gtg aaa aag aag acc caa ggg aag gtt gta	685
Ala Arg Ile Asn Ser His Val Lys Lys Thr Gln Gly Lys Val Val	
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Asp Ile Ile Gln Gly Leu Asp Leu Thr Ala Met Val Leu Val Asn	
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His Ile Phe Phe Lys Ala Lys Trp Glu Lys Pro Phe His Leu Glu Tyr	
225 230 235	
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Thr Arg Lys Asn Phe Pro Phe Leu Val Gly Glu Gln Val Thr Val Gln	
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Val Pro Met Met His Gln Lys Glu Gln Phe Ala Phe Gly Val Asp Thr	
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Glu Leu Asn Cys Phe Val Leu Gln Met Asp Tyr Lys Gly Asp Ala Val	
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Ala Phe Phe Val Leu Pro Ser Lys Gly Lys Met Arg Gln Leu Glu Gln	
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gcc ttg tca gcc aga aca ctg ata aag tgg agc cac tca ctc cag aaa	1021
Ala Leu Ser Ala Arg Thr Leu Ile Lys Trp Ser His Ser Leu Gln Lys	
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 Arg Trp Ile Glu Val Phe Ile Pro Arg Phe Ser Ile Ser Ala Ser Tyr  
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 Asn Leu Glu Thr Ile Leu Pro Lys Met Gly Ile Gln Asn Ala Phe Asp  
 335 340 345 350

aaa aat gct gat ttt tct gga att gca aag aga gac tcc ctg cag gtt 1165  
 Lys Asn Ala Asp Phe Ser Gly Ile Ala Lys Arg Asp Ser Leu Gln Val  
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tct aaa gca acc cac aag gct gtg ctg gat gtc agt gaa gag gcc act 1213  
 Ser Lys Ala Thr His Lys Ala Val Leu Asp Val Ser Glu Glu Gly Thr  
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gag gcc aca gca gct acc acc acc aag ttc ata gtc cga tgg aag gat 1261  
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 Gly Pro Ser Tyr Phe Thr Val Ser Phe Asn Arg Thr Phe Leu Met Met  
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 Asn Pro Thr Lys Ser  
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Cys Ala Pro Ile Tyr Cys Val Ser Pro Ala Asn Ala Pro Ser Ala Tyr  
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Pro Arg Pro Ser Ser Thr Lys Ser Thr Pro Ala Ser Gln Val Tyr Ser  
50 55 60

Leu Asn Thr Asp Phe Ala Phe Arg Leu Tyr Arg Arg Leu Val Leu Glu  
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Thr Pro Ser Gln Asn Ile Phe Phe Ser Pro Val Ser Val Ser Thr Ser  
85 90 95

Leu Ala Met Leu Ser Leu Gly Ala His Ser Val Thr Lys Thr Gln Ile  
100 105 110

Leu Gln Gly Leu Gly Phe Asn Leu Thr His Thr Pro Glu Ser Ala Ile  
115 120 125

His Gln Gly Phe Gln His Leu Val His Ser Leu Thr Val Pro Ser Lys  
130 135 140

Asp Leu Thr Leu Lys Met Gly Ser Ala Leu Phe Val Lys Lys Glu Leu  
145 150 155 160

Gln Leu Gln Ala Asn Phe Leu Gly Asn Val Lys Arg Leu Tyr Glu Ala  
165 170 175

Glu Val Phe Ser Thr Asp Phe Ser Asn Pro Ser Ile Ala Gln Ala Arg  
180 185 190

Ile Asn Ser His Val Lys Lys Lys Thr Gln Gly Lys Val Val Asp Ile  
195 200 205

Ile Gln Gly Leu Asp Leu Leu Thr Ala Met Val Leu Val Asn His Ile  
210 215 220

Phe Phe Lys Ala Lys Trp Glu Lys Pro Phe His Leu Glu Tyr Thr Arg  
225 230 235 240

Lys Asn Phe Pro Phe Leu Val Gly Glu Gln Val Thr Val Gln Val Pro  
245 250 255

Met Met His Gln Lys Glu Gln Phe Ala Phe Gly Val Asp Thr Glu Leu  
260 265 270

Asn Cys Phe Val Leu Gln Met Asp Tyr Lys Gly Asp Ala Val Ala Phe  
275 280 285

Phe Val Leu Pro Ser Lys Gly Lys Met Arg Gln Leu Glu Gln Ala Leu  
290 295 300

Ser Ala Arg Thr Leu Ile Lys Trp Ser His Ser Leu Gln Lys Arg Trp  
305 310 315 320

Ile Glu Val Phe Ile Pro Arg Phe Ser Ile Ser Ala Ser Tyr Asn Leu  
325 330 335

Glu Thr Ile Leu Pro Lys Met Gly Ile Gln Asn Ala Phe Asp Lys Asn  
340 345 350

Ala Asp Phe Ser Gly Ile Ala Lys Arg Asp Ser Leu Gln Val Ser Lys  
355 360 365

Ala Thr His Lys Ala Val Leu Asp Val Ser Glu Glu Gly Thr Glu Ala  
370 375 380

Thr Ala Ala Thr Thr Thr Lys Phe Ile Val Arg Ser Lys Asp Gly Pro  
385 390 395 400

Ser Tyr Phe Thr Val Ser Phe Asn Arg Thr Phe Leu Met Met Ile Thr  
405 410 415

Asn Lys Ala Thr Asp Gly Ile Leu Phe Leu Gly Lys Val Glu Asn Pro  
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Thr Lys Ser  
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Asn Leu Thr Gln Glu Asn Gln Asp Arg Gly Thr His Val Asp Leu Gly  
35 40 45

Leu Ala Ser Ala Asn Val Asp Phe Ala Phe Ser Leu Tyr Lys Gln Leu  
50 55 60

Val Leu Lys Ala Pro Asp Lys Asn Val Ile Phe Ser Pro Leu Ser Ile  
65 70 75 80

Ser Thr Ala Leu Ala Phe Leu Ser Leu Gly Ala His Asn Thr Thr Leu  
85 90 95

Thr Glu Ile Leu Lys Gly Leu Lys Phe Asn Leu Thr Glu Thr Ser Glu  
100 105 110

Ala Glu Ile His Gln Ser Phe Gln His Leu Leu Arg Thr Leu Asn Gln  
115 120 125

Ser Ser Asp Glu Leu Gln Leu Ser Met Gly Asn Ala Met Phe Val Lys  
130 135 140

Glu Gln Leu Ser Leu Leu Asp Arg Phe Thr Glu Asp Ala Lys Arg Leu  
145 150 155 160

Tyr Gly Ser Glu Ala Phe Ala Thr Asp Phe Gln Asp Ser Ala Ala Ala  
165 170 175

Lys Lys Leu Ile Asn Asp Tyr Val Lys Asn Gly Thr Arg Gly Lys Ile  
180 185 190

Thr Asp Leu Ile Lys Asp Leu Asp Ser Gln Thr Met Met Val Leu Val  
195 200 205

Asn Tyr Ile Phe Phe Lys Ala Lys Trp Glu Met Pro Phe Asp Pro Gln  
210 215 220

Asp Thr His Gln Ser Arg Phe Tyr Leu Ser Lys Lys Lys Trp Val Met  
225 230 235 240

Val Pro Met Met Ser Leu His His Leu Thr Ile Pro Tyr Phe Arg Asp  
245 250 255

Glu Glu Leu Ser Cys Thr Val Val Glu Leu Lys Tyr Thr Gly Asn Ala  
260 265 270

Ser Ala Leu Phe Ile Leu Pro Asp Gln Asp Lys Met Glu Glu Val Glu  
275 280 285

Ala Met Leu Leu Pro Glu Thr Leu Lys Arg Trp Arg Asp Ser Leu Glu  
290 295 300

Phe Arg Glu Ile Gly Glu Leu Tyr Leu Pro Lys Phe Ser Ile Ser Arg  
305 310 315 320

Asp Tyr Asn Leu Asn Asp Ile Leu Leu Gln Leu Gly Ile Glu Glu Ala  
325 330 335

Phe Thr Ser Lys Ala Asp Leu Ser Gly Ile Thr Gly Ala Arg Asn Leu  
340 345 350

Ala Val Ser Gln Val Val His Lys Ala Val Leu Asp Val Phe Glu Glu  
355 360 365

Gly Thr Glu Ala Ser Ala Ala Thr Ala Val Lys Ile Thr Leu Leu Ser  
370 375 380

Ala Leu Val Glu Thr Arg Thr Ile Val Arg Phe Asn Arg Pro Phe Leu  
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Met Ile Ile Val Pro Thr Asp Thr Gln Asn Ile Phe Phe Met Ser Lys  
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Val Thr Asn Pro Lys Gln Ala  
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35 40 45

Leu Lys Ile Ala Pro Ala Asn Ala Asp Phe Ala Phe Arg Phe Tyr Tyr  
50 55 60

Leu Ile Ala Ser Glu Thr Pro Gly Lys Asn Ile Phe Phe Ser Pro Leu  
65 70 75 80

Ser Ile Ser Ala Ala Tyr Ala Met Leu Ser Leu Gly Ala Cys Ser His  
85 90 95

Ser Arg Ser Gln Ile Leu Glu Gly Leu Gly Phe Asn Leu Thr Glu Leu  
100 105 110

Ser Glu Ser Asp Val His Arg Gly Phe Gln His Leu Leu His Thr Leu  
115 120 125

Asn Leu Pro Gly His Gly Leu Glu Thr Arg Val Gly Ser Ala Leu Phe  
130 135 140

Leu Ser His Asn Leu Lys Phe Leu Ala Lys Phe Leu Asn Asp Thr Met  
145 150 155 160

Ala Val Tyr Glu Ala Lys Leu Phe His Thr Asn Phe Tyr Asp Thr Val  
165 170 175

Gly Thr Ile Gln Leu Ile Asn Asp His Val Lys Lys Glu Thr Arg Gly  
180 185 190

Lys Ile Val Asp Leu Val Ser Glu Leu Lys Lys Asp Val Leu Met Val  
195 200 205

Leu Val Asn Tyr Ile Tyr Phe Lys Ala Leu Trp Glu Lys Pro Phe Ile  
210 215 220

Ser Ser Arg Thr Thr Pro Lys Asp Phe Tyr Val Asp Glu Asn Thr Thr  
225 230 235 240

Val Arg Val Pro Met Met Leu Gln Asp Gln Glu His His Trp Tyr Leu  
245 250 255

His Asp Arg Tyr Leu Pro Cys Ser Val Leu Arg Met Asp Tyr Lys Gly  
260 265 270

Asp Ala Thr Val Phe Phe Ile Leu Pro Asn Gln Gly Lys Met Arg Glu  
275 280 285

Ile Glu Glu Val Leu Thr Pro Glu Met Leu Met Arg Trp Asn Asn Leu  
290 295 300

Leu Arg Lys Arg Asn Phe Tyr Lys Lys Leu Glu Leu His Leu Pro Lys  
305 310 315 320

Phe Ser Ile Ser Gly Ser Tyr Val Leu Asp Gln Ile Leu Pro Arg Leu  
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Gly Phe Thr Asp Leu Phe Ser Lys Trp Ala Asp Leu Ser Gly Ile Thr  
340 345 350

Lys Gln Gln Lys Leu Glu Ala Ser Lys Ser Phe His Lys Ala Thr Leu  
355 360 365

Asp Val Asp Glu Ala Gly Thr Glu Ala Ala Ala Ala Thr Thr Phe Ala  
370 375 380

Ile Lys Phe Phe Ser Ala Gln Thr Asn Arg His Ile Leu Arg Phe Asn  
385 390 395 400

Arg Pro Phe Leu Val Val Ile Phe Ser Thr Ser Thr Gln Ser Val Leu  
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Phe Leu Gly Lys Val Val Asp Pro Thr Lys Pro  
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Thr	Ile	His	Cys	Ala	Ser	Pro	Glu	Gly	Lys	Val	Thr	Ala	Cys	His	Ser
		20						25					30		
Ser	Gln	Pro	Asn	Ala	Thr	Leu	Tyr	Lys	Met	Ser	Ser	Ile	Asn	Ala	Asp
	35							40					45		
Phe	Ala	Phe	Asn	Leu	Tyr	Arg	Arg	Phe	Thr	Val	Glu	Thr	Pro	Asp	Lys
	50					55					60				
Asn	Ile	Phe	Phe	Ser	Pro	Val	Ser	Ile	Ser	Ala	Ala	Leu	Val	Met	Leu
65				70						75					80
Ser	Phe	Gly	Ala	Cys	Cys	Ser	Thr	Gln	Thr	Glu	Ile	Val	Glu	Thr	Leu
			85						90					95	
Gly	Phe	Asn	Leu	Thr	Asp	Thr	Pro	Met	Val	Glu	Ile	Gln	His	Gly	Phe
		100					105						110		
Gln	His	Leu	Ile	Cys	Ser	Leu	Asn	Phe	Pro	Lys	Lys	Glu	Leu	Glu	Leu
		115					120					125			
Gln	Ile	Gly	Asn	Ala	Leu	Phe	Ile	Gly	Lys	His	Leu	Lys	Pro	Leu	Ala
	130					135					140				
Lys	Phe	Leu	Asn	Asp	Val	Lys	Thr	Leu	Tyr	Glu	Thr	Glu	Val	Phe	Ser
145				150						155					160
Thr	Asp	Phe	Ser	Asn	Ile	Ser	Ala	Ala	Lys	Gln	Glu	Ile	Asn	Ser	His
			165						170					175	
Val	Glu	Met	Gln	Thr	Lys	Gly	Lys	Val	Val	Gly	Leu	Ile	Gln	Asp	Leu
		180						185					190		
Lys	Pro	Asn	Thr	Ile	Met	Val	Leu	Val	Asn	Tyr	Ile	His	Phe	Lys	Ala
	195						200					205			
Gln	Trp	Ala	Asn	Pro	Phe	Asp	Pro	Ser	Lys	Thr	Glu	Asp	Ser	Ser	Ser
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Phe	Leu	Ile	Asp	Lys	Thr	Thr	Thr	Val	Gln	Val	Pro	Met	Met	His	Gln
225				230					235					240	
Met	Glu	Gln	Tyr	Tyr	His	Leu	Val	Asp	Met	Glu	Leu	Asn	Cys	Thr	Val
			245						250					255	
Leu	Gln	Met	Asp	Tyr	Ser	Lys	Asn	Ala	Leu	Ala	Leu	Phe	Val	Leu	Pro
		260						265					270		
Lys	Glu	Gly	Gln	Met	Glu	Ser	Val	Glu	Ala	Ala	Met	Ser	Ser	Lys	Thr

275

280

285

Leu Lys Lys Trp Asn Arg Leu Leu Gln Lys Gly Trp Val Asp Leu Phe  
290 295 300

Val Pro Lys Phe Ser Ile Ser Ala Thr Tyr Asp Leu Gly Ala Thr Leu  
305 310 315 320

Leu Lys Met Gly Ile Gln His Ala Tyr Ser Glu Asn Ala Asp Phe Ser  
325 330 335

Gly Leu Thr Glu Asp Asn Gly Leu Lys Leu Ser Asn Ala Ala His Lys  
340 345 350

Ala Val Leu His Ile Gly Glu Lys Gly Thr Glu Ala Ala Val Pro  
355 360 365

Glu Val Glu Leu Ser Asp Gln Pro Glu Asn Thr Phe Leu His Pro Ile  
370 375 380

Ile Gln Ile Asp Arg Ser Phe Met Leu Leu Ile Leu Glu Arg Ser Thr  
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Arg Ser Ile Leu Phe Leu Gly Lys Val Val Asn Pro Thr Glu Ala  
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35 40 45

Asn Ile Phe Phe Ser Pro Val Ser Ile Ala Thr Ala Phe Ala Met Leu  
50 55 60

Ser Leu Gly Thr Lys Ala Asp Thr His Asp Glu Ile Leu Glu Gly Leu  
65 70 75 80

Asn Phe Asn Leu Thr Glu Ile Pro Glu Ala Gln Ile His Glu Gly Phe  
85 90 95

Gln Glu Leu Leu Arg Thr Leu Asn Gln Pro Asp Ser Gln Leu Gln Leu  
100 105 110

Thr Thr Gly Asn Gly Leu Phe Leu Ser Glu Gly Leu Lys Leu Val Asp  
115 120 125

Lys Phe Leu Glu Asp Val Lys Lys Leu Tyr His Ser Glu Ala Phe Thr  
130 135 140

Val Asn Phe Gly Asp Thr Glu Glu Ala Lys Lys Gln Ile Asn Asp Tyr  
145 150 155 160

Val Glu Lys Gly Thr Gln Gly Lys Ile Val Asp Leu Val Lys Glu Leu  
165 170 175

Asp Arg Asp Thr Val Phe Ala Leu Val Asn Tyr Ile Phe Phe Lys Gly  
180 185 190

Lys Trp Glu Arg Pro Phe Glu Val Lys Asp Thr Glu Glu Asp Phe  
195 200 205

His Val Asp Gln Val Thr Thr Val Lys Val Pro Met Met Lys Arg Leu  
210 215 220

Gly Met Phe Asn Ile Gln His Cys Lys Lys Leu Ser Ser Trp Val Leu  
225 230 235 240

Leu Met Lys Tyr Leu Gly Asn Ala Thr Ala Ile Phe Phe Leu Pro Asp  
245 250 255

Glu Gly Lys Leu Gln His Leu Glu Asn Glu Leu Thr His Asp Ile Ile  
260 265 270

Thr Lys Phe Leu Glu Asn Glu Asp Arg Arg Ser Ala Ser Leu His Leu  
275 280 285

Pro Lys Leu Ser Ile Thr Gly Thr Tyr Asp Leu Lys Ser Val Leu Gly  
290 295 300

Gln Leu Gly Ile Thr Lys Val Phe Ser Asn Gly Ala Asp Leu Ser Gly  
305 310 315 320

Val Thr Glu Glu Ala Pro Leu Lys Leu Ser Lys Ala Val His Lys Ala  
325 330 335

Val Leu Thr Ile Asp Glu Lys Gly Thr Glu Ala Ala Gly Ala Met Phe  
340 345 350

Leu Glu Ala Ile Pro Met Ser Ile Pro Pro Glu Val Lys Phe Asn Lys  
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Pro Phe Val Phe Leu Met Ile Glu Gln Asn Thr Lys Ser Pro Leu Phe  
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Met Gly Lys Val Val Asn Pro Thr Gln Lys  
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Gln Val Ser Lys Ala Thr His Lys Ala Val Leu Asp Val  
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Leu Glu

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Thr

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